STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

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THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.4.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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 U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street,
 Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER:			
ATTN: NEW RULES CASES:	PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE			
Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."			
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.			
3Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters , instead.			
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.			
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.			
-	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.			
(OLD RULĖS)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped			
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.			
•	Sequence(s) missing. If intentional , please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000			
(NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.			
Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence			
11 Use of <220>	Sequence(e) missing the <220> "Feature" and associated numeric identifiers and responses.			
2/	Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)			
"bug"	Please do not use "Copy to Disk" function of Patentln version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.			
13 Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid			



IFWP

RAW SEQUENCE LISTING DATE: 03/15/2007 PATENT APPLICATION: US/10/558,155 TIME: 11:30:32

Input Set: A:\2007-01-03 1254-0299PUS1.txt
Output Set: N:\CRF4\03152007\J558155.raw

3 <110> APPLICANT: WAKITA, Takaji KATO, Takanobu 4 DATE, Tomoko <120> TITLE OF INVENTION: A NUCLEIC ACID CONSTRUCT CONTAINING A NUCLEIC ACID DERIVED FROM THE GENOME OF HEPATITIS C VIRUS (HCV) OF GENOTYPE 2a, 9 AND A CELL HAVING SUCH NUCLEIC ACID CONSTRUCT INTRODUCED THEREIN 11 <130> FILE REFERENCE: 1254-0299PUS1 13 <140> CURRENT APPLICATION NUMBER: US 10/558,155 14 <141> CURRENT FILING DATE: 2005-11-23 16 <150> PRIOR APPLICATION NUMBER: PCT/JP2003/015038 17 <151> PRIOR FILING DATE: 2003-11-25 Dees Not Comply 19 <150> PRIOR APPLICATION NUMBER: JP 2003-148242 20 <151> PRIOR FILING DATE: 2003-05-26 Corrocted Diskette Nex 22 <150> PRIOR APPLICATION NUMBER: JP 2003-329115 23 <151> PRIOR FILING DATE: 2003-09-19 15 explain source of ofgenetic material 25 <160> NUMBER OF SEO ID NOS: 41 27 <170> SOFTWARE: PatentIn Ver. 2.1 29 <210> SEQ ID NO: 1 30 <211> LENGTH: 8024 31 <212> TYPE: RNA 32 <213> ORGANISM: Artificial Sequence 34 <220> FEATURE: 35 <223> OTHER INFORMATION: Description of Artificial Sequence: replicon 37 <400> SEQUENCE: 1 38 accugeeccu aauaggggeg acaeucegee augaaucaeu eeccugugag gaacuaeugu 60 39 cuucacgcag aaagcgccua gccauggcgu uaguaugagu gucguacagc cuccaggccc 120 40 cccccucccg ggagagccau aguggucugc ggaaccggug aguacaccgg aauugccggg 180 41 aagacugggu ccuuucuugg auaaacccac ucuaugcccg gccauuuggg cgugcccccg 240 42 caagacugcu agccgaguag cguuggguug cgaaaggccu ugugguacug ccugauaggg 300 43 cgcuugcgag ugccccggga ggucucguag accgugcacc augagcacaa auccuaaacc 360 44 ucaaagaaaa accaaaagaa acaccaaccg ucgcccaaug auugaacaag auggauugca 420 45 cgcagguucu ccggccgcuu ggguggagag gcuauucggc uaugacuggg cacaacagac 480 46 aaucggeuge ueugaugeeg eeguguueeg geugueageg eaggggegee egguueuuuu 540 47 ugucaagacc gaccuguccg gugcccugaa ugaacugcag gacgaggcag cgcggcuauc 600 48 guggcuggcc acgacgggcg uuccuugcgc agcugugcuc gacguuguca cugaagcggg 660 49 aagggacugg cugcuauugg gcgaagugcc ggggcaggau cuccugucau cucaccuugc 720 50 uccugeegag aaaguaueea ucauggeuga ugeaaugegg eggeugeaua egeuugauee 780 51 ggcuaccugc ccauucgacc accaagcgaa acaucgcauc gagcgagcac guacucggau 840 52 ggaagccggu cuugucgauc aggaugaucu ggacgaagag caucaggggc ucgcgccagc 900 53 cgaacuguuc gccaggcuca aggcgcgcau gcccgacggc gaggaucucg ucgugaccca 960 54 uggcgaugec ugcuugecga auaucauggu ggaaaaugge egcuuuucug gauucaucga 1020 55 cuguggccgg cugggugugg cggaccgcua ucaggacaua gcguuggcua cccgugauau 1080 56 ugcugaagag cuuggcggcg aaugggcuga ccgcuuccuc gugcuuuacg guaucgccgc 1140

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Output Set: N:\CRF4\03152007\J558155.raw

57 ucccgauucg cagcgcaucg ccuucuaucg ccuucuugac gaguucuucu gaguuuaaac 1200 58 ccucucccuc cccccccu aacguuacug gccgaagccg cuuggaauaa ggccggugug 1260 59 cguuugucua uauguuauuu uccaccauau ugccgucuuu uggcaaugug agggcccgga 1320 60 aaccuggeee ugucuucuug acgageauuc cuaggggucu uuccecucuc gecaaaggaa 1380 61 ugcaaggucu guugaauguc gugaaggaag caguuccucu ggaagcuucu ugaagacaaa 1440 62 caacgucugu agcgacccuu ugcaggcagc ggaacccccc accuggcgac aggugccucu 1500 63 geggecaaaa gecaegugua uaagauacae eugeaaagge ggeacaaeee cagugecaeg 1560 64 uugugaguug gauaguugug gaaagaguca aauggcucuc cucaagcgua uucaacaagg 1620 65 ggcugaagga ugcccagaag guaccccauu guaugggauc ugaucugggg ccucggugca 1680 66 caugcuuuac auguguuuag ucgagguuaa aaaaacgucu aggccccccg aaccacgggg 1740 67 acgugguuuu ccuuugaaaa acacgaugau accauggcuc ccaucacugc uuaugcccag 1800 68 caaacacgag gccuccuggg cgccauagug gugaguauga cggggcguga caggacagaa 1860 69 caggeegggg aagueeaaau ceugueeaca gueucueagu eeuueeuegg aacaaceaue 1920 70 ucggggguuu uguggacugu uuaccacgga gcuggcaaca agacucuagc cggcuuacgg 1980 71 gguccgguca cgcagaugua cucgagugcu gagggggacu ugguaggcug gcccagcccc 2040 72 ccugggacca agucuuugga gccgugcaag uguggagccg ucgaccuaua ucuggucacg 2100 73 cggaacgcug augucauccc ggcucggaga cgcggggaca agcggggagc auugcucucc 2160 74 cegagaccca uuucgaccuu gaaggggucc ucgggggggc cggugcucug cccuaggggc 2220 75 cacqueguug ggcucuuceg ageageugug ugcucueggg geguggeeaa auceauegau 2280 76 uucauccccg uugagacacu cgacguuguu acaaggucuc ccacuuucag ugacaacagc 2340 77 acgccaccgg cugugcccca gaccuaucag gucggguacu ugcaugcucc aacuggcagu 2400 78 ggaaagagca ccaagguccc ugucgcguau gccgcccagg gguacaaagu acuagugcuu 2460 79 aaccceuegg uageugecae ceugggguuu ggggeguaee uauceaagge acauggeaue 2520 80 aaucccaaca uuaggacugg agucaggacc gugaugaccg gggaggccau cacguacucc 2580 81 acauauggca aauuucucgc cgaugggggc ugcgcuaggg gcgccuauga caucaucaua 2640 82 ugcgaugaau gccacgcugu ggaugcuacc uccauucucg gcaucggaac gguccuugau 2700 83 caagcagaga cagccggggu cagacuaacu gugcuggcua cggccacacc ccccggguca 2760 84 gugacaaccc cccaucccga uauagaagag guaggccucg ggcgggaggg ugagaucccc 2820 85 uucuauggga gggcgauucc ccuauccugc aucaagggag ggagacaccu gauuuucugc 2880 86 cacucaaaga aaaaguguga cgagcucgcg gcggcccuuc ggggcauggg cuugaaugcc 2940 87 guggcauacu auagaggguu ggacgucucc auaauaccag cucagggaga uguggugguc 3000 88 guegecaceg aegeceueau gaegggguae aeuggagaeu uugaeueegu gauegaeuge 3060 89 aauguagegg ucaeccaage uguegaeuue ageeuggaee ceaecuucae uauaaecaea 3120 90 cagacugucc cacaagacgc ugucucacgc agucagcgcc gcgggcgcac agguagagga 3180 91 agacagggca cuuauaggua uguuuccacu ggugaacgag ccucaggaau guuugacagu 3240 92 guagugcuuu gugagugcua cgacgcaggg gcugcguggu acgaucucac accagcggag 3300 93 accaccguca ggcuuagagc guauuucaac acgcccggcc uacccgugug ucaagaccau 3360 94 cuugaauuuu gggaggcagu uuucacegge cucacacaca uagaegeeca cuuccucuc 3420 95 caaacaaagc aagcggggga gaacuucgcg uaccuaguag ccuaccaagc uacggugugc 3480 96 gccagagcca aggccccucc cccguccugg gacgccaugu ggaagugccu ggcccgacuc 3540 97 aagccuacgc uugcgggccc cacaccucuc cuguaccguu ugggcccuau uaccaaugag 3600 98 gucacccuca cacacccugg gacgaaguac aucgccacau gcaugcaagc ugaccuugag 3660 99 gucaugacca gcacgugggu ccuagcugga ggaguccugg cagccgucgc cgcauauugc 3720 100 cuggcgacug gaugcguuuc caucaucggc cgcuugcacg ucaaccagcg agucgucguu 3780 101 gcgccggaua aggagguccu guaugaggcu uuugaugaga uggaggaaug cgccucuagg 3840 102 gcggcucuca ucgaagaggg gcagcggaua gccgagaugu ugaaguccaa gauccaaggc 3900 103 uugcugcagc aggccucuaa gcaggcccag gacauacaac ccgcuaugca ggcuucaugg 3960 104 cccaaagugg aacaauuuug ggccagacac auguggaacu ucauuagcgg cauccaauac 4020 105 cucgcaggau ugucaacacu gccagggaac cccgcggugg cuuccaugau ggcauucagu 4080

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			cauaggccug				
			ggcccucguc				
110	ucuauggaag	augucaucaa	ucuacugccu	gggauccugu	cuccgggagc	ccugguggug	4380
111	ggggucaucu	gcgcggccau	ucugcgccgc	cacgugggac	cgggggaggg	cgcgguccaa	4440
			cuuugcuucc				
113	gugacggagu	cggaugcguc	gcagcgugug	acccaacuac	uuggcucucu	uacuauaacc	4560
114	agccuacuca	gaagacucca	caauuggaua	acugaggacu	gccccauccc	augcuccgga	4620
			ggacuggguu				
			caagcugccc				
			cacuggcauc				
			gggcucuaug				
			uaucaauugc				
			caucuggagg				
			uguaacagga				
			uuucuccugg				
			ccgggaugag				
			cugugaaccu				
			cacggcggag				
			cucagugagc				
			cuaugacgug				
			gccugagucc				
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			auacuccugg				
			caacccuuug				
			gagcgccuca				
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			caccuuggag				
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154	ggcgaugacc	uaguagucau	cucagaaagc	caggggacug	aggaggacga	gcggaaccug	7020

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155 agagecuuca eggaggecau gaecagguae ueugeceeue euggugauee eeceagaeeg 7080 156 gaauaugacc uggagcuaau aacauccugu uccucaaaug ugucuguggc guugggcccg 7140 157 eggggeegee geagauaeua eeugaeeaga gaeeeaaeea euceaeuege eegggeugee 7200 158 ugggaaacag uuagacacuc cccuaucaau ucauggcugg gaaacaucau ccaguaugcu 7260 159 ccaaccauau ggguucgcau gguccuaaug acacacuucu ucuccauucu caugguccaa 7320 160 gacacccugg accagaaccu caacuuugag auguauggau caguauacuc cgugaauccu 7380 161 uuggaccuuc cagccauaau ugagagguua cacgggcuug acgccuuuuc uaugcacaca 7440 162 uacucucace acgaacugae geggguggeu ucageceuca gaaaacuugg ggegeeaeee 7500 163 cucagggugu ggaagagucg ggcucgcgca gucagggcgu cccucaucuc ccguggaggg 7560 164 aaagcggccg uuugcggccg auaucucuuc aauugggcgg ugaagaccaa gcucaaacuc 7620 165 acuccauuge eggaggegeg ecuacuggae uuauecaguu gguucaeegu eggegeegge 7680 166 gggggcgaca uuuuucacag cgugucgcgc gcccgacccc gcucauuacu cuucggccua 7740 167 cuccuacuuu ucguaggggu aggecucuuc cuacuccceg cucgguagag cggcacacac 7800 169 uuuuuuuuu cuuuuuuuu uuuuucccuc uuucuucccu ucucaucuua uucuacuuuc 7920 170 uuucuuggug geuccaucuu ageecuague aeggeuageu gugaaaggue egugageege 7980 171 augacugcag agagugccgu aacuggucuc ucugcagauc augu 174 <210> SEQ ID NO: 2 175 <211> LENGTH: 8024 See item See item # 11 on error surmary surmary 176 <212> TYPE: RNA 177 <213> ORGANISM: Artificial Sequence 179 <220> FEATURE: 180 <223> OTHER INFORMATION: Description of Artificial Sequence: {replicon 182 <400> SEQUENCE: 2 183 accegececu aauaggggeg acaeucegee augaaucaeu eeeeugugag gaacuaebgu 60 184 cuucacgcag aaagcgucua gccauggcgu uaguaugagu gucguacagc cuccaggccc 120 185 cccccucceg ggagagccau aguggucugc ggaaccggug aguacaccgg aauugccggg 180 186 aagacugggu ccuuucuugg auaaacccac ucuaugcccg gccauuuggg cgugcccccg 240 187 caagacugcu agccgaguag cguuggguug cgaaaggccu ugugguacug ccugauaggg 300 188 ugcuugcgag ugccccggga ggucucguag accgugcacc augagcacaa aucccaaacc 360 189 ucaaagaaaa accaaaagaa acacuaaccg ucgcccaaug auugaaçaag auggauugca 420 190 cgcagguucu ccggccgcuu ggguggagag gcuauucggc uaugacuggg cacaacagac 480 191 aaucggcugc ucugaugccg ccguguuccg gcugucagcg caggggcgcc cgguucuuuu 540 192 ugucaagacc gaccuguccg gugcccugaa ugaacugcag gacgaggcag cgcggcuauc 600 193 guggcuggcc acgacgggcg uuccuugcgc agcugugcuc gacguuguca cugaagcggg 660 194 aagggacugg cugcuauugg gcgaagugcc ggggcaggau cuccugucau cucaccuugc 720 195 uccugeegag aaaguaucea ucauggeuga ugeaaugegg eggeugeaua egeuugauee 780 196 ggcuaccugc ccauucgacc accaagcgaa acaucgcauc gagcgagcac guacucggau 840 197 ggaagceggu cuugucgauc aggaugaucu ggacgaagag caucaggggc ucgcgccagc 900 198 cgaacuguuc gccaggcuca aggcgcgcau gcccgacggc gaggaucucg ucgugaccca 960 199 uggcgaugcc ugcuugccga auaucauggu ggaaaauggc cgcuuuucug gauucaucga 1020 200 cuguggcegg cugggugugg cggaccgcua ucaggacaua gcguuggcua cccgugauau 1080 201 ugcugaagag cuuggcggcg aaugggcuga ccgcuuccuc gugcuuuacg guaucgccgc 1140 202 uccegauucg cagegeaucg ceuucuaucg ceuucuugac gaguucuucu gaguuuaaac 1200 203 ccucucccuc ccccccccu aacguuacug gccgaagccg cuuggaauaa ggccggugug 1260 204 eguuugucua uauguuauuu uccaccauau ugccgucuuu uggcaaugug agggcccgga 1320 205 aaccuggeee ugucuucuug acgageauue cuaggggueu uuccecucuc gecaaaggaa 1380 206 ugcaaggucu guugaauguc gugaaggaag caguuccucu ggaagcuucu ugaagacaaa 1440 207 caacgucugu agcgacccuu ugcaggcagc ggaacccccc accuggcgac aggugccucu 1500-The type of errors shown exist throughout

the Begilence Listing. Please check subsequent

Estilences for similar orrors.

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Input Set : A:\2007-01-03 1254-0299PUS1.txt
Output Set: N:\CRF4\03152007\J558155.raw

208 gcggccaaaa gccacgugua uaagauacac cugcaaaggc ggcacaaccc cagugccacg 1560 209 uugugaguug gauaguugug gaaagaguca aauggcucuc cucaagcgua uucaacaagg 1620 210 ggcugaagga ugcccagaag guaccccauu guaugggauc ugaucugggg ccucggugca 1680 211 caugcuuuac auguguuuag ucgagguuaa aaaaacgucu aggccccccg aaccacgggg 1740 212 acgugguuuu ccuuugaaaa acacgauaau accauggccc ccaucaccgc uuacgcccag 1800 213 cagacacgag gucucuuggg cucuauagug gugagcauga cggggcguga caagacagaa 1860 214 caggccgggg agguccaagu ccuguccaca gucacucagu ccuuccucgg aacauccauu 1920 215 ucgggggucu uauggacugu uuaccacgga gcuggcaaca agacacuagc cggcucgcgg 1980 216 ggcccgguca cgcagaugua cucgagcgcc gagggggacu uggucgggug gcccagcccu 2040 217 ccugggacca aaucuuugga gccguguacg uguggagcgg ucgaccugua uuuggucacg 2100 218 cggaacgcug augucauccc ggcucgaaga cgcggggaca agcggggagc gcugcucucc 2160 219 ccgagacccc uuucgaccuu gaaggggucc ucggggggac cugugcuuug cccuaggggc 2220 220 cacgcugucg gaaucuuccg ggcagcugug ugcucucggg guguggcuaa guccauagau 2280 221 uucauccccg uugagacgcu cgacaucguc acgcggucuc ccaccuuuag ugacaacagc 2340 222 acaccaccag cugugececa gaccuaucag gugggguacu ugcacgecec cacuggeagu 2400 223 ggaaaaagca ccaagguccc cgucgcguac gccgcccagg gguauaaagu gcuggugcuc 2460 224 aaucccucgg uggcugccac ccugggauuu ggggcguacu uguccaaggc acauggcauc 2520 225 aaccccaaca uuaggacugg agucagaacu gugacgaccg gggagcccau uacauacucc 2580 226 acguauggua aauuccucgc cgaugggggc ugcgcaggcg gcgccuauga caucaucaua 2640 227 ugcgaugaau gccacucugu ggaugcuacc acuauucucg gcaucgggac aguccuugac 2700 228 caagcagaga cagccggggu caggcuaacu guacuggcca cggccacgcc ccccgggucg 2760 229 gugacaaccc cccaucccaa uauagaggag guagcccucg gacaggaggg ugagaucccc 2820 230 uucuauggga gggcguuucc ccugucuuac aucaagggag ggaggcacuu gauuuucugc 2880 231 cacucaaaga aaaaguguga cgagcucgca acggcccuuc ggggcauggg cuugaacgcu 2940 232 guggcauauu acagaggguu ggacgucucc auaauaccaa cucaaggaga uguggugguc 3000 233 guugccaccg acgcccucau gacgggguau acuggagacu uugacuccgu gaucgacugc 3060 234 aacguagegg ucacccagge eguagacuuc agecuggace ecaccuucae uauaaccaca 3120 235 cagacuguce egeaagaege ugucucaegu agucagegee gagggegeae ggguagagga 3180 236 agacugggca uuuauaggua uguuuccacu ggugagcgag ccucaggaau guuugacagu 3240 237 guaguacucu gugagugcua cgacgcagga gcugcuuggu augagcucuc accaguggag 3300 238 acgaccguca ggcucagggc guauuucaac acgccuggcu ugccugugug ccaggaccac 3360 240 cagacaaagc agucggggga aaauuucgca uacuuaguag ccuaucaggc cacagugugc 3480 242 aagcccacgc uugugggccc uacaccucuc cuguaccguu ugggcucugu uaccaacgag 3600 243 gucacccuua cacaccccgu gacaaaauac aucgccacau gcaugcaagc ugaccucgag 3660 244 gucaugacca gcacgugggu ccuggcuggg ggagucuuag cagccgucgc cgcguauugc 3720 245 uuagegaeeg gguguguuue caucauugge eguuuaeaea ueaaeeageg ageuguegue 3780 246 gcuccggaca aggagguccu cuaugaggcu uuugaugaga uggaggaaug ugccuccaga 3840 247 gcggcucucc uugaagaggg gcagcggaua gccgagaugc ugaaguccaa gauccaaggc 3900 248 uuauugcage aageeucuaa acaggeecag gacauacaae eegeugugea ageuuegugg 3960 249 cccaagaugg agcaauucug ggccaaacau auguggaacu ucauaagcgg cauucaguac 4020 250 cucgcaggac ugucaacacu gccagggaac ccugcugugg cuuccaugau ggcauucagc 4080 251 gccgcccuca ccaguccguu gucaacuagc accaccaucc uucuuaacau ucuggggggc 4140 252 uggcuggegu cccaaauugc gccacccgcg ggggccacug gcuuuguugu caguggccug 4200 253 gugggagcug cuguuggcag cauaggcuug gguaaagugc ugguggacau ccuggcaggg 4260 254 uauggugegg gcauuueggg ggcceuegue geguuuaaga ucaugueugg egagaageee 4320 255 uccauggagg augucaucaa cuugcugccu gggauucugu cuccaggugc ucugguggug 4380 256 ggagucaucu gcgcggccau ucugcgccgc caugugggac cgggggaagg cgcgguccaa 4440 VERIFICATION SUMMARY

DATE: 03/15/2007 TIME: 11:30:33

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PATENT APPLICATION: US/10/558,155